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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
10/810,733	03/26/2004	Qiong Cheng	CL2385USNA	3038
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E I DU PONT DE NEMOURS AND COMPANY			LEE, JAE W	
LEGAL PATENT RECORDS CENTER BARLEY MILL PLAZA 25/1128		ART UNIT	PAPER NUMBER	
4417 LANCASTER PIKE WILMINGTON, DE 19805			1656	
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Please find below and/or attached an Office communication concerning this application or proceeding.

		Application No.	Applicant(s)		
Office Action Summary		10/810,733	CHENG ET AL.		
		Examiner	Art Unit		
		Jae W. Lee	1656		
Period fo	The MAILING DATE of this communication apport	pears on the cover sheet with the c	orrespondence address		
A SH WHI(- Exte after - If NO - Failu Any	ORTENED STATUTORY PERIOD FOR REPLICATION OF THE MAILING DON'S INC. SIX (6) MONTHS from the mailing date of this communication. Operiod for reply is specified above, the maximum statutory period vire to reply within the set or extended period for reply will, by statute reply received by the Office later than three months after the mailing ed patent term adjustment. See 37 CFR 1.704(b).	ATE OF THIS COMMUNICATION 36(a). In no event, however, may a reply be tim will apply and will expire SIX (6) MONTHS from 6, cause the application to become ABANDONE	N. nely filed the mailing date of this communication. D (35 U.S.C. § 133).		
Status					
1)⊠ 2a)⊠ 3)□	Responsive to communication(s) filed on Of This action is FINAL. 2b) This Since this application is in condition for alloware closed in accordance with the practice under Expression 1.	s action is non-final. nce except for formal matters, pro			
Disposit	ion of Claims				
 4) ☐ Claim(s) 15,17-19,36 and 37 is/are pending in the application. 4a) Of the above claim(s) is/are withdrawn from consideration. 5) ☐ Claim(s) 36 is/are allowed. 6) ☐ Claim(s) 15,17-19 and 37 is/are rejected. 7) ☐ Claim(s) is/are objected to. 8) ☐ Claim(s) are subject to restriction and/or election requirement. 					
Applicat	ion Papers				
10)□	The specification is objected to by the Examine The drawing(s) filed on is/are: a) acc Applicant may not request that any objection to the Replacement drawing sheet(s) including the correct The oath or declaration is objected to by the Ex	epted or b) objected to by the Idea of a by the Idea of a by the Idea of the I	e 37 CFR 1.85(a). jected to. See 37 CFR 1.121(d).		
Priority (under 35 U.S.C. § 119				
 12) Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f). a) All b) Some * c) None of: 1. Certified copies of the priority documents have been received. 2. Certified copies of the priority documents have been received in Application No 3. Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)). * See the attached detailed Office action for a list of the certified copies not received. 					
2) 🔲 Notic	t(s) e of References Cited (PTO-892) e of Draftsperson's Patent Drawing Review (PTO-948) nation Disclosure Statement(s) (PTO/SB/08)	4) ☐ Interview Summary Paper No(s)/Mail Da 5) ☐ Notice of Informal P	ate		
	r No(s)/Mail Date	6) Other:			

Application/Control Number: 10/810,733

Art Unit: 1656

DETAILED ACTION

Application status

Applicants' amendment to claims 6 (cancelled), 18 (amended) and 19 (amended), and the addition of new claims 36 and 37, filed on 09/07/2006, is acknowledged. Applicants' arguments, filed on 09/07/2006, are acknowledged. Claims 15, 17-19, 36 and 37 are at issue and present for examination. Applicants' arguments filed on 09/07/2006, have been fully considered, and are deemed to be persuasive to overcome some of the rejections previously applied. Rejections and/or objections not reiterated from previous office actions are hereby withdrawn.

Election/Restrictions

Applicants' traversed the restriction requirement for claims 4, 5, and 20-35. However, claims 4, and 5 are drawn to a polypeptide, classified in class 435, subclass 183; claims 20-27 are drawn to a method of producing carotenoid compounds, classified in class 435, subclass 67; claims 28-32 are drawn to a method of regulating carotenoid biosynthesis in an organism, classified in class 435, subclass 471; claim 33 is drawn to a strain DC413, classified in class 435, subclass 252.3; and claim 34 is drawn to all isolated nucleic acid molecule encoding all of the amino acid sequences as set forth in SEQ ID NO: 2, 4, 6, 8, 10, 12, and 14, classified in class 536, subclass 23.1. Because these inventions are classified differently and they are patentably distinct for the reasons stated in the non-final rejection, filed on 08/30/2006, Applicants' arguments

is found non-persuasive and the restriction requirement is deemed proper under 35 U.S.C. 121.

Claim Objections

Claim 35 is objected to because it depends from a non-existent claim.

Claims 15 and 17-19 are objected to because they depend from the cancelled claim 6, which is necessitated by the Applicants' amendment, filed on 09/07/2006.

Appropriate correction is required.

Claim Rejections - 35 USC § 112

The following is a quotation of the first paragraph of 35 U.S.C. 112:

The specification shall contain a written description of the invention, and of the manner and process of making and using it, in such full, clear, concise, and exact terms as to enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make and use the same and shall set forth the best mode contemplated by the inventor of carrying out his invention.

Claims 15, 17-19 and 37 are rejected under 35 U.S.C. 112, first paragraph, because the specification, while being enabling for an isolated nucleic acid molecule of SEQ ID NO: 20 encoding all of the following enzymes a) geranylgeranyl pyrophosphate synthetase of SEQ ID NO: 2; b) zeaxanthine glucosyl transferase of SEQ ID NO: 4; c) lycopene cyclase of SEQ ID NO: 6; d) phytoene desaturase of SEQ ID NO: 8; e) phytoene synthase of SEQ ID NO: 10; f) β-carotene hydroxylase of SEQ ID NO: 12; and g) isopentenyl diphosphate isomerase of SEQ ID NO: 14., does not reasonably provide enablement for any isolated nucleic acid having 95% identity to SEQ ID NO 20 and

encoding all of the following enzymes (a)-(g). The specification does not enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make and use the invention commensurate in scope with these claims.

Factors to be considered in determining whether undue experimentation is required are summarized in In re Wands (858 F.2d 731, 8 USPQ 2nd 1400 (Fed. Cir. 1988)) as follows: (1) the quantity of experimentation necessary, (2) the amount of direction or guidance presented, (3) the presence or absence of working examples, (4) the nature of the invention, (5) the state of the prior art, (6) the relative skill of those in the art, (7) the predictability or unpredictability of the art, and (8) the breadth of the claim(s).

Claim 37 is so broad as to encompass any nucleic acid having 95% sequence identity to SEQ ID NO 20 encoding all of the following enzymes a) any geranylgeranyl pyrophosphate synthetase; b) any zeaxanthine glucosyl transferase; c) any lycopene cyclase; d) any phytoene desaturase; e) any phytoene synthase; f) any β-carotene hydroxylase; and g) any isopentenyl diphosphate isomerase (see above 112 2nd paragraph rejection). The scope of the claims is not commensurate with the enablement provided by the disclosure with regard to the extremely large number of nucleic acids broadly encompassed by the claims, encoding the polypeptides, a) geranylgeranyl pyrophosphate synthetase; b) zeaxanthine glucosyl transferase; c) lycopene cyclase; d) phytoene desaturase; e) phytoene synthase; f) β-carotene hydroxylase; and g) isopentenyl diphosphate isomerase, by any nucleic acid having 95% identity to SEQ ID NO 20. The claims rejected under this section of U.S.C. 112,

first paragraph, place no structure/function limitations on the claimed nucleic acids.

Since the amino acid sequence of a protein determines its structural and functional

properties, predictability of which changes can be tolerated in a protein's amino acid

sequence and thus nucleic acid sequence while obtaining the desired activity requires a

knowledge of and guidance with regard to which amino acids in the protein's sequence,

if any, are tolerant of modification and which are conserved (i.e. expectedly intolerant to

modification), and detailed knowledge of the ways in which the protein's structure

relates to its function. While recombinant and mutagenesis techniques are known, it is

not routine in the art to screen for multiple substitutions or multiple modifications among

any of the 9127 nucleic acid residues or any 456 nucleic acid residues of SEQ ID NO:

20 (5 % of the SEQ ID NO: 20), as encompassed by the instant claims, and the

positions within a protein's sequence where amino acid modifications can be made with

a reasonable expectation of success in obtaining the desired activity/utility are limited in

any protein and the result of such modifications is unpredictable. In addition, one skilled

in the art would expect any tolerance to modification for a given protein to diminish with

each further and additional modification, e.g. multiple substitutions.

It is noted that claims 15 and 17-19 continue to be rejected to the extent that it is Applicants' intent that they depend from claim 37, which was newly added for the purpose of replacing claim 6, as disclosed in the Applicants' argument on pg. 9, line 1-2, filed on 09/07/2006. Thus, said claims are continued to be rejected because they improperly depend from the cancelled claim 6.

of preferred codons for specific host cell expression) may be made to such a gene cluster without affecting its utility provided that all of the encoded polypeptides are expressed and are enzymatically active. Applicants have not provided guidance to make a majority of the encompassed isolated nucleic acid molecule having at least 95% identity to SEQ ID NO 20, wherein said nucleic acid molecule encodes all of the enzymes a) geranylgeranyl pyrophosphate synthetase; b) zeaxanthine glucosyl transferase; c) lycopene cyclase; d) phytoene desaturase; e) phytoene synthase; f) βcarotene hydroxylase; and g) isopentenyl diphosphate isomerase. The specification does not support the broad scope of the claims because the specification does not establish: (A) regions of the nucleic acid structure which may be modified without affecting the encoded proteins' functions/activities; (B) the general tolerance of the protein to modification and the extent of such tolerance; (C) a rational and predictable scheme for modifying SEQ ID NO: 20 thereby changing amino acid residue of the encoded polypeptides with an expectation of obtaining the desired biological function; and (D) the insufficient guidance in the specification as to which of the essentially infinite possible choices is likely to be successful. Because of this lack of guidance, the extended experimentation that would be required to determine which substitutions at which location among as many as 456 nucleotides (5% of the SEQ ID NO 20), would be acceptable to retain the desired polypeptide functions/activities and the fact that the relationship between the sequence of a peptide and its tertiary structure (i.e. its activity) are not well understood and are not predictable (e.g., see Ngo et al. in The Protein Folding Problem and Tertiary Structure Prediction, 1994, Merz et al. (ed.), Birkhauser,

Boston, MA, pp. 433 and 492-495, Ref: U, Form-892), it would require undue experimentation for one skilled in the art to arrive at the majority of those nucleic acid molecules.

Thus, applicants have not provided sufficient guidance to enable one of ordinary skill in the art to make and use the claimed invention in a manner reasonably correlated with the scope of the claims broadly including any isolated nucleic acid molecule having at least 95% sequence identity to SEQ ID NO 20 and encoding all of the following enzymes: a) geranylgeranyl pyrophosphate synthetase; b) zeaxanthine glucosyl transferase; c) lycopene cyclase; d) phytoene desaturase; e) phytoene synthase; f) β-carotene hydroxylase; and g) isopentenyl diphosphate isomerase. The scope of the claims must bear a reasonable correlation with the scope of enablement (In re Fisher, 166 USPQ 19 24 (CCPA 1970)). Without sufficient guidance, determination of those nucleic acids having the desired biological characteristics is unpredictable and the experimentation left to those skilled in the art is unnecessarily, and improperly, extensive and undue. See In re Wands 858 F.2d 731, 8 USPQ2nd 1400 (Fed. Cir, 1988).

Although, Applicants argue that the number of nucleic acid molecules that encode all the recited polypeptide having 95% identity to SEQ ID NO: 20 is very small while retaining the enzymatic function of each protein, a skilled artisan would not recognize which 456 nucleotides out of 9127 would can be substituted, modified or changed and still maintain desired level of enzymatic activity for each protein.

Therefore, there is still insufficient guidance to practice the invention with respect to this issue.

Further, it is noted that Applicants have amended claim 18 in response to a rejection under this statue for the inclusion of green plants. However, claims 17-19 still encompass a transformed host from green plants. Previous studies demonstrate that the transformation of plants or the gene transfer to plants have been difficult with low success rate (Potrykus, Biotechnology 8(6): 535-542). In this regard, the specification fails to disclose a representative example of a successful transformation of a green plant host with the nucleic acid molecule of SEQ ID NO 20. Thus, Applicants have not provided sufficient guidance to make such.

Remarks

Claims 15, 17-19 and 37 are not allowed.

Conclusion

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Jae W. Lee whose telephone number is 571-272-9949. The examiner can normally be reached on 8:30 am - 5:00 pm.

If attempts to reach the examiner by telephone are unsuccessful, please contact the examiner's supervisor, Kathleen Kerr at 571-272-0931. The fax phone number for the organization where this application or proceeding is assigned is 571-273-9949.

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atent Examiner: Jae W. Lee, Ph.D.

RICHARD HUTSON, PH.D. PRIMARY EXAMINER